

## SEQUENCE LISTING



<110> ORCHID BIOSCIENCES, INC.  
 MCKEOWN, Brian  
 DERBYSHIRE, Roger  
 ROWAN, Paul  
 SUNG, Robert

<120> Methods and Compositions for Genotyping

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<140> 10/632,393  
 <141> 2003-08-01

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 <151> 2002-08-02

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 <151> 2002-12-20

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ataaccaaaa gggctaagtg gaagagacat ataattactt attctaaaat tgtagaattt 480  
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&lt;222&gt; (231)...(231)

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agtgycatga gcaggcctct tatacathtt ggcaatgact atgaggaccg ttactatcrt 180
gaaaacatgt accgttaccc caaccaagtg tactacagac cagtggatcr ktatagtaac 240
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accaccaagg                                     310

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&lt;211&gt; 420

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&lt;213&gt; Ovis aries

&lt;400&gt; 44

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gagctgctgc agctggagca gtggtagggg gccttgggtgg ctacatgctg ggaagtggca 180
tgagcaggcc tcttatacat tttggcaatg actatgagga ccgttactat cgtgaaaaca 240
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actttgtgca tgactgtgtc aacatcacag tcaagcaaca cacagtcacc accaccacca 360
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&lt;211&gt; 192

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gctgtcctgg aa                                     192

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&lt;211&gt; 422

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&lt;213&gt; Ovis aries

&lt;400&gt; 46

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tttcccacac aggggtatatt accttaggtg tcctaagaga tttttgcctt tcaatgtacg 360
caagcccagc acatgccatg gtacatagta gagattttct ctctctcgct gtctctctct 420
ct                                     422

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&lt;211&gt; 226

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&lt;213&gt; Ovis aries

&lt;400&gt; 47

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ccagggtggac tgctttttct cccgcttaca tttcctagaa agtgccctgc tcaccctttt 180
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13428US.TXT

SEQUENCE LISTING

<110> BIOSCIENCES, INC.  
MCKEOWN, Brian  
DERBYSHIRE, Roger  
ROWAN, Paul  
SUNG, Robert

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tgccagacca aaggggtagg gaccgtactc tagagaccct caccacacag gatgattaaa 600
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ataacccaaa gggctaagtg gaagagacat ataattactt attctaaaat tgtagaattt 480  
ggcaagaagt gagacttatt tgattcattt ataaaacatg taaacaaaag acagttatcc 540  
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<210> 36  
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<400> 36  
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 <223> r = a or g  
  
 <220>  
 <221> misc\_feature  
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 <223> r = a or g  
  
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 <221> misc\_feature

<222> (125)...(125)  
 <223> y, = t or c

<220>  
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 <222> (231)...(231)  
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 agtgycatga gcaggcctct tatacathtt ggcaatgact atgaggaccg ttactatcrt 180  
 gaaaacatgt accgttacc ccaaccaagt tactacagac cagtggatcr ktatagtaac 240  
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 accaccaagg 310

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 gagctgctgc agctggagca gtggtagggg gccttgggtgg ctacatgctg ggaagtggca 180  
 tgagcaggcc tcttatacat tttggcaatg actatgagga ccgttactat cgtgaaaaca 240  
 tgtaccgtta cccaaccaa gtgtactaca gaccagtggg tgggtatagt aaccagaaca 300  
 actttgtgca tgactgtgtc aacatcacag tcaagcaaca cacagtcacc accaccacca 360  
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 <212> DNA  
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 gctgtcctgg aa 192

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 ggagatctat gcccttctcc aagttctggg gaagaagggt tggagacagc cactacccaa 180  
 aatgtathtt tcttcattct tcaccttgct aaatcttacg cattttaagg tcccagcttt 240  
 ctattcctcc attcaaaaaa cataatttga ttatttgctt ctattccaat attctttgta 300  
 tttccacac agggtatatt accttaggtg tcttaagaga tttttgcctt tcaatgtacg 360  
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 ct 422

<210> 47  
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 <212> DNA  
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<400> 47  
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 ccagggtggac tgctttttct cccgcttaca tttcctagaa agtgccctgc tcaccctttt 180  
 ctctggatgc tcaactcagg gttttaccag gcctgaactc tctcca 226